===
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: markspencer
Timestamp: [year=2009; month=11; day=18; hr=14; min=30; sec=11; ms=788;
]
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Reviewer Comments:
1.
E320   Wrong Nucleic Acid Designator, gc in SEQID (481)

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<213> Homo sapiens
<220>
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Val Thr Lys Ser Phe Asn Lys Gly Glu Cys Lys Gly Glu Phe Ala
    210   215   220
<210> 482
<211> 223
<212> PRT

Codons must be made up of three nucleotides. The last codon in SEQ ID #
481 is only two nucleotides. Please make all necessary changes.

2.
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W213   Artificial or Unknown found in <213> in SEQ ID (2)
W213   Artificial or Unknown found in <213> in SEQ ID (3)

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W213            Artificial or Unknown found in <213> in SEQ ID (6)
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W213            Artificial or Unknown found in <213> in SEQ ID (15)
W213            Artificial or Unknown found in <213> in SEQ ID (16)
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W213            Artificial or Unknown found in <213> in SEQ ID (18)
W213            Artificial or Unknown found in <213> in SEQ ID (19)
W213            Artificial or Unknown found in <213> in SEQ ID (20)  This
error has occured more than 20 times, will not be displayed
E257            Invalid sequence data feature in <221> in SEQ ID (636)
E257            Invalid sequence data feature in <221> in SEQ ID (636)
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The warnings and errors shown above are ok and require no response.
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Application No: 10045674   Version No: 4.0

Input Set:

Output Set:

Started: 2009-11-05 17:45:21.376

Finished: 2009-11-05 17:45:29.384

Elapsed: 0 hr(s) 0 min(s) 8 sec(s) 8 ms

Total Warnings: 459

Total Errors: 9

No. of SeqIDs Defined: 637

Actual SeqID Count: 637

Error code	Error Description
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W 213	Artificial or Unknown found in <213> in SEQ ID (19)
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            Input Set:



            Output Set:




                Started:   2009-11-05 17:45:21.376
               Finished:   2009-11-05 17:45:29.384
                Elapsed:   0 hr(s) 0 min(s) 8 sec(s) 8 ms
        Total Warnings:    459
          Total Errors:    9
  No. of SeqIDs Defined:   637
      Actual SeqID Count:  637
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Error code	Error Description
	This error has occured more than 20 times, will not be displayed
E 320	Wrong Nucleic Acid Designator, gc in SEQID (481)
E 257	Invalid sequence data feature in <221> in SEQ ID (636)
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<110> LADNER, ROBERT C.
      COHEN, EDWARD H.
      NASTRI, HORACIO G.
      ROOKEY, KRISTIN L.
      HOET, RENE
      HOOGENBOOM, HENDRICUS R. J. M.

<120> NOVEL METHODS OF CONSTRUCTING LIBRARIES COMPRISING
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      OF PEPTIDES, POLYPEPTIDES OR PROTEINS AND THE NOVEL
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<141> 2001-10-25

<150> 06/198,069
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<150> 09/837,306
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<160> 637

<170> PatentIn Ver. 3.5

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